

PCT09

RAW SEQUENCE LISTING DATE: 01/31/2003 PATENT APPLICATION: US/09/743,414A TIME: 11:52:10

Input Set : A:\NIELSEN4.txt

Output Set: N:\CRF4\01312003\I743414A.raw

```
3 <1100 APPLICANT: Nielsen, Jens
              Nissen, Torben L
              Hielland-Brandt, Morten C
        	imes 1.00 \cdot TITLE 0F INVENTION: Metabolically engineered microbial cell with an altered
              metabolite production
     10 <1 * ← FILE REFERENCE: NIELSEN#4
     11 4140 - CUREENT APPLICATION NUMBER: 09/743,414A
     15 <141 - CURRENT FILING DATE: 2001-01-10
     15 :150 - PRICE APPLICATION NUMBER: PCT/DK99/00397
     16 <1:1: PRICE FILING DATE: 1994-07-12
     18 R1 TO BRICE APPLICATION NUMBER: PA 1995 00967
     14 (1:1 - PAICA FILING DATE: 1993-07-10
     Al <160 - NUMBER OF SEQ ID NOS: 14
     As <170× COFTWARE: Patentin Ver. 2.1
     15 KO160 PEQ ID NO: 1
     20 KLII: MENGTH: 1395
     37 K. TIDA TYPE: DMA
     가 로 IR + ORGANISM: Azotobacter vinelandii
     FEATURE:
     31 KINTA NAME/KEY: CDS
     St K. 122 LOCATION: (1)..(1395)
     45 KLIBA OTHER INFORMATION:
   > 35 <400> 1
     He after dos gits that was tab gat gitg gitg gits alto ggo wow ggo cot got
                                                                              48
W--> 37 Met Ala Val Tyr Xaa Tyr Asp Val Val Val Ile Gly Thr Gly Pro Ala
                                                                               96
     40 gặc daw gọc ghá gog átg ast dọc gtg làng góc ggg các aag gta gọc
     41 Gly Gly Gly Ala Ala Met Ash Ala Val Lys Ala Gly Arg Lys Val Ala
                     20°
     1.
     44 yra ata gat gat egu cee cag ate ggo ago aac tgo aco cac oto gga
                                                                               144
     15 Val Val Asp Asp Arg Pro Glm Val Gly Gly Asm Cys Thr His Leu Gly
                                                                               1.32
     4F ann att edd ted aag gog otg ogd han tog gig ogg dag ald atg dag
     4 + Thr Ile Pro Ser Lys Ala Leu Arg His Ser Val Arg Gln Ile Met Gln
             50
                                                                              240
      I had aan aac aat dig dig the ego dag atd ggd gag dog ogs igg itt
      is Tyr Ash Ash Ash Pro Leu Phe Arg Gln Ile Gly Glu Pro Arg Trp Pho
                              7 ÷
     to to: the god gat give etg aag age god gag dag give ate god aag dag
                                                                              288
        Se: Phe Ala Asp Val leu Lys Ser Ala Glu Gln Val Ile Ala Lys Gln
                          50
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     61 Va. Ser Ser Arg Thr Gly Tyr Tyr Ala Arg Ash Arg Ile Asp Thr Phe
```



DATE: 01 31/2003

RAW SEQUENCE LISTING

FATERT AFFILICATION: US/09/743,414A TIME: 11:82:11

Input Set : A:\NIELSEN4.txt

Output Set: N:\CRF4\01312003\I743414A.raw

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		Ph€	Gly		Ala	Ser	Phe	СУЗ		Glu	His	Thr	Ile		Val	Val	His	
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	901 9		210	7.7.7	2 5 7	a 5.4	5 5.3	215 ata	~* .~	at.a	,7,7,7	030	220	or a ri	a	tac	17.11.17	720
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		225					235					235				- 1	243	
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RAW SEQUENCE LISTING

DATE: 01/31/2003 FATENT APPLICATION: US/09/743,414A TIME: 11:52:10

Input Set : A:\NIELSEN4.txt

Output Set: N:\CRF4\01312003\I743414A.raw

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1z8 gft root tac gag gto ggo sag gor the threship ggr ang groung gra-
  129 Val Pro Tyr Glu Val Gly Lys Ala Phe Phe Lys Gly Met Ala Arg Ala
                3 15
  170 370
                                   390
  1. Hag at higher give gag sag give ggd atgletg sag sith old bit cachego.
                                                                          1200
  1:4 Gln II. Ala Val Glu Lys Ala Gly Met Lei Lys Ile Leu Phe His Arg
                           390
                                              395
  1\% , agracy storigae atolete gge gtd casinge the ggs hat day get teg
                                                                          1248
  1.4 Glu Thr Leu Glu Ile Leu Gly Val His Cys Phe Gly Tyr Gln Ala Ser
                                           410
                      405
  14) yaw ato gio bat aid ggo bag gog aid aig aad bag aag ggo gag gog
  141 Glu II: Val His lle Gly Gln Ala Ile Met Asn Gir Lys Gly Glu Ala
  14:
                  4.00
                                                           430
                                       4.25
  144 aut ach ath aug tat the ate and acc acc the auc tac deg acc atg
                                                                          1344
  14% Akh Thr Dev Lys Tyr Phe Ile Ash Thr Thr Phe Ash Tyr Pro Thr Met
                                  4.4\,0
                                                      4.45
  14\% quantizating gain two logicity graphics gain gain gat attempt aat legg att tit.
                                                                          1392
  14 + Ala Glo Ala Tyr Ara Val Ala Ala Tyr Asp Gly Lea Asm Ara Leu Phe
  15
                              455
                                                   4.60
  15. Squ
                                                                          1395
  156 - . 10 - . EQ 10 NO: 1
  15 - ... II - DENGTH: 464
  119 HILL TYPE: PRT
  100 -013 - OEGANISM: Anotobacter vinelandii
  1.1 SLIGH FEATURE:
  101 HIII HAME/KEY: misc feature
  163 + 0000 + bocation: (5)...(5)
  164 GLLS OTHER INFORMATION: The 'Mas' at location 5 stands for Lys, or Asn.
  166 S. . C FEATURE:
  10% HLL1: NAME/KEY: misc feature
  108 -0.01 - LOCATION: (304)..(304)
  164 CLCS - OTHER INFORMATION: The 'Maa' at location 304 stands for Lys, or Asn.
  171 - 400 - SEQUENCE: 3
-> 172 Met Ala Val Tyr Xaa Tyr Asp Val Val Val Ile Gly Thr Gly Pro Ala
  1%5 Gly Glu Gly Ala Ala Met Asn Ala Val Lys Ala Gly Arg Lys Val Ala
                  . 0
                                        2.5
  100 Val Val Asp Asp Arg Pro Glr Val Gly Gly Ash Cys Thr His Leu Gly
  174
              3.57
                                   40
                                                        4.5
```

1%: Thr : The Pro Ser Lys Ala Leu Ard His Ser Val Ard Gln : The Met Gln 4.5 180 50 184 Tyr Asr. Asr. Akr. Pro Leu Phe Arg Gln Ilc Gly Glu Pro Arg Trp Phe 7 C 75 185 187 Fer Phe Ala Asp Val Leu Lys Ser Ala Glu Gln Val Ile Alā Lys Gln 188 85 90 1.00 Val. Ser Ser Arg Thr Gly Tyr Tyr Ala Arg Asn Arg Ile Asp Thr Phe 100 1 11 105 110 103 File Gly Thr Ala Ser Phe Cys Asp Glu His Thr Ile Giu Val Val His 120 196 Leu Asn Gly Met Val Glu Thr Leu Val Ala Lys Gln Phe Val Ile Ala



PATENT APPLICATION: US/09/743,414A

DATE: 01/31/2003 TIME: 11:52:55

Input Set : A:\NIELSEN4.txt

Output Set: N:\CRF4\01312003\I743414A.raw

```
199 Thr Gly Ser Arg Pro Tyr Arg Fro Ala Asp Val Asp Phe Thr His Pro
  3mr 145 150
                                           155
  29. Arg lie Tyr Asp Ser Asp Thr lie Leu Ser Leu Gly His Thr Fro Arg
  1.45
                                                           175
                                     1.70
  🗈 E Arg Leu Ile Ile Tyr Gly Ala Gly Val Ile Gly Cyg Glu Tyr Ala Ser
                                    185
  1888 Ile Phe Ser Gly Leu Gly Val Leu Val Asp Leu Ile Asp Asn Arg Asp
  .594 195
                                 () ()
                                                    265
  Mi Gin Leu Leu Ser Phe Ser Asp Asp Glu Ile Ser Asp Ser Leu Ser Tyr
  .11. 216
                             .215
                                                221
  ::14 His Let Arg Ash Ash Ash Val Beu Ile Arg His Ash Glu Glu Tyr Glu
                                            230
                        2 30
  .17 Arg Val Glu Gry Leu Asp Asr Gly Val Ile Leo His Leu Lys Ser Gly
                     2 4 5
                                        250
  Lit lays bys life bys Ara Asp Ala Pho Leu Trp Ser Asr. G.y Ang Thr Gly
                 26
                                    165
  ... Ach Thr Asp Dys Deu Gly Deu Glu Ash Ile Gly Leu Lys Ala Ash Gly
  275
                                                285
                            2.84
-> 226 Arg Gly Gln Ile Gln Val Asp Glu His Tyr Arg Thr Glu Val Ser Xaa
  ...7 291
                            2.95
  ... tire Tyr Ala Ala Gly Asp Val Ilo Gly Trp Pro Ser Lou Ala Ser Ala
  , 30 305
                                            315
                         54 O
  . I Ala Tyr Asp Glm Gly Ard Sor Ala Ala Gly Ser Ile Thr Glu Ash Asp
                     3. 5
                                        5. 3 Ú
  .:5 Car Trp Arg Phe Val Asp Asp Val Pro Thr Gly Ile Tyr Thr 1le Pro
                 340
                                     345
                                                        350
  . A- Glu lie Ser Ser Val Gly Lys Thr Glu Arg Glu Leu Thr Glm Ala Dys
  154
             355
                                 :61
                                                    3+.5
  .41 Val Pro Tyr Giu Val Gly Lys Ala Phe Phe Lyz Gly Met Ala Arg Ala
                             375
  144 G.n The Ala Val Glu by: Ala Gly Met Leu by: The Leu Phe His Arg
                                            3, 9 5
  .47 Glu Thr Leu Glu lie Leu Gly Val His Cys Pho Gly Tyr Glr Ala Ser
                     415
                                        410
  220 Glu Ile Val His Ile Gly Gln Ala Ile Met Ash Gin Lys Gly Glu Ala
                 4.30
                                    4.25
                                                       430
  ich Ash Thr Leu Lys Tyr Phe Ile Ash Thr Thr Phe Ash Tyr Pro Thr Met
  . 4455
                                440
                                        4.4.5
   Ne Ala Gly Ala Tyr Arg Vai Ala Ala Tyr Asp Gly Leu Asn Arg Leu Phe
  150
                             455
  , 61 -0.10 × SEQ ID NO: ≥
  . CT -1.11 - LENGTH: 52
  . + + + 1. 12 ≥ TYPE: DNA
  . 대 전 13 * ORGANISM: Artificial Sequence
  . +++ -1.20> FEATURE:
  1.67 (123) OTHER INFORMATION: Description of Artificial Sequence: Primer
  164 -1400> SEQUENCE: 3
  270 arteategat gaattetate thatggtees attetthact geachgitha ca
```

DATE: 01/31/2003

TIME: 11:52:10

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/743,414A

Imput Net : A:\NIELSEN4.txt

Output Set: N:\CRF4\01312003\I743414A.raw

273 - 2162 SEO ID NO: 4 274 <2112 LEMGTH: 11 275 <212> TYPE: DN4 ..76 72134 OR MANISM: Artificial Sequence .78 <220 - FEMATURE: 279 (22) (OTHER INFORMATION: Description of Artificial Sequence: Frimer 281 Stort SEQUENCE: 4 UNL pp://wet.ast/datatewaaag/cattefeteg/et.ggttaatt/tteetgtete/ttgtetatea/60 353 quantamaa a 2*6 Killer SED ID NO: 5 PAR BURIE DEMIGTH: 24 298 -COLOR TYPE: DNA 199 1017 ORBANISM: Artificial Sequence .W. C.Bus FRATURE: . 4. OTHER INFORMATION: Description of Artificial Sequence: Primer .94 F400 - SENULTICE: 5 . " grantan is aat diccaaa agad 2.4 . We of Los SEQ III NO: 6 .9+ 4.11 - LENGTH: 34 WO E. L. TYPE: DNA # 1 F. 17 - OEFANISM: Artificial Sequence FOR HOLDER STATUMES: 204 C. 200 OTHER INFORMATION: Description of Artificial Sequence: Primer afoo -dioo - £EQUENCE: 6 2007 grag ragato fictagaatg offittigata acaaaaat 3.8 310 d. 10 - SEQ 10 NO: 7 311 d. 11 - DENGTH: 38 Fil Hull - TYPE: ONA olr of loo OH WANISM: Artificial Sequence Sie e. Gererature: Fig. 4. A: OTHER INFORMATION: Description of Artificial Sequence: Primer G18 -MOOR SEQUENCE: 7 619 brodolagato teogelogaga geetaalaega ttaacaaa 58 377 - 116 - SE2 IL NO: 8 BAY THIN ELENGTH: 19 304 K. ILM TYPE: DNA 3.5 - 11 - OEGANISM: Artificial Sequence FIG. 4. LOS FEATURE: Fig. 4...2 OTHER INFORMATION: Description of Artificial Sequence: Primer 330 MADON SEPTEMBE: 8 Mil greamacase aaggiceta 19 334 -0010. SP₂ 1D NC: 9 Addoor: The LENGTH: 38 630 CHILD TYPE: DNA 237 - 2130 OFGANISM: Artificial Sequence 239 - 22200 FEATURE: 340 <22230 CTHER INFORMATION: Description of Artificial Sequence: Frimer 341 <400> SEQUENCE: 9 345 gogogggato ototaquatg coagtgttga aatoagad 3.8

RAW SEQUENCE LISTING ERROR SUMMARY PATENT APPLICATION: US/09/743,414A

DATE: %1/31/2003 TIME: 11:52:11

Input Cet : A:\NIELSEN4.txt

Output Set: N:\CRF4\01312003\1743414A.raw

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:1; Xaa Pos. 5,304
Seq#:2; Xaa Pos. 5,304

VERIFICATION SUMMARY

PATENT APPLICATION: US/09/743,414A

DATE: 01/31/2003 TIME: 11:52:11

Input Set : A:\NIELSEN4.txt

Output Set: N:\CRF4\01312003\I743414A.raw

L:35 M:258 W: Mandatory Feature missing, <2232 Blank for CEQ#:1, bine#:33 L:37 M:341 W: (46) "n" or "Xaa" used, for SEQ 1D#:1 after pos.:48 L:109 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:912 L:172 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:0 L:226 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:288